

TABLE III

MOTIFS	POSITION					
	1° anchor 1	2	3	4	5	6
DR4 preferred deleterious	FMYLIVW	M	T	W	I	VSTCPALIM MH R
DR1 preferred deleterious	MFLIVWY	C	CH	PAMQ FD	CWD	VMATSPILIC M GDE
DR7 preferred deleterious	MFLIVWY	M	W	A	G	IVMSACTPL M GRD
DR Supermotif	MFLIVWY					VMSTACPLI IV G
DR3 MOTIFS	1° anchor 1	2	3	1° anchor 4	5	1° anchor 6
motif a preferred	LIVMFY			D		
motif b preferred	LIVMFAY			DNQEST		KRH

Italicized residues indicate less preferred or "tolerated" residues.

SP 1028-1

(SEQ ID NO: 2457)  
(SEQ ID NO: 2436)

***Amendments***

***In the Specification:***

B1 Please cancel the existing Sequence Listing for the above-identified application, replace it with the substitute Sequence Listing appended hereto, and insert the same at the end of the application.

B2 Please replace pending page 99 of the specification with the attached page 99 of the specification.